



OIPE

RAW SEQUENCE LISTING

4 <110> APPLICANT: Jones, Chris L.

PATENT APPLICATION: US/09/766,880B

DATE: 03/21/2003 TIME: 13:38:06

Input Set : A:\110163-132US2.ST25.txt
Intput Set: N:\CRF4\03212003\I766880B.raw

```
Abrams, Ezra S.
             Eleffer-Higgins, Stephen G.
              Thang, Tianhong
             Eacock, Graham P.
             Well, Machael
             McLowell, Christopher S.
     11 - 120 - TITLE OF INVENTION: Methods of and Apparatus for Separating and Detecting
Nuclein Acad
     14 - (180) - FILE PEFERENCE: 110163.132US2
     16 - 1400 CUFRENT APPLICATION NUMBER: US 09/766,880B
     1 - - 1411 - CUPPENT FILING DATE: 2001-01-19
                                                              ENTERED
     1 + -15 - PHIOF APPLICATION NUMBER: US 60/176,839
     10 (181) PRIOR FILING DATE: 2001-01-19
     THE RIGHT NOSE SEQ ID NOSE 20
     14 (17) COUNTRE: FastSEQ for Windows Version 4.0
     DO RELEASED IN NO. 1
     17 - 1201 / LENGIH: 50
     Da K. 12 & TYPE: EMA
     .3 -0.13 - OFFAMISM: Staphylococcus epidermis
     31 /400. SEQUENCE: 1
                                                                             50
     30 matimassat of saggiost gasggaagsa gsattaagtg gatestsata.
     34 + 310 + SEQ ID NO: 3
     35 KU11 - LENGTH: 36
     36 KULD - TYFE: DNA
     37 - 1313 - ORGANISM: Unknown
     33 KLEGO FEATURE:
     40 km25 - OTHER INFORMATION: staphylococcus
     42 -: 400 - SEQUENCE: 2
                                                                             26
     4% acatytosyg teotgabyga agbago
     45 -010 - SEQ ID NO: 3
     46 -1211 - LENGTE: 30
     47 -110 - TYFE: DNA
     48 -011 - OKGANISM: Staphylococcus aureus
     34 -400 - BEQUENCE: 3
                                                                             51)
     11 Sattaastat of raggioot gaoggaagoa goattaagtg gatoatoata
     13 SHIN - SHE ID NO: 4
     ta Kulli - LENGTH: 49
     55 HOLD - TYPE: DNA
     FW +001 + + ORGANISM: Unknown
     th Hidder FEATURE:
     ( ) - 0.223 - OTHER INFORMATION: staphylococcus
     61 (400 / SEQUENCE: 4
                                                                             49
     62 catyaaceat gtoaggtoot gaoggaagea geattaagtg gatotoata
```

RAW SEQUENCE LISTING DATE: 03/21/2003 PATENT APPLICATION: US/09/766,880B TIME: 13:58:06

Input Set : A:\110163-132US2.ST25.txt Output Set: N:\CRF4\03212003\I766880B.raw

64 <210> SEO ID NO: 5	
65 (211)> LENGTH: 46	
on C21.15 TYPE: DUE	
하는 401일 - ORMANEUM: Streptococcus pyogenes	
NOTES OF SEQUENCES: 5	
The officing might maken right da liggiggaggaat libbagbageed staagee	46
- N. HAMI - PERMITH MO: 6	
10 - 1211 - DENGTH: 40	
14 - 1211 - TYPE: 1004	
11 -0.11 : OF MANI.TI: Streptococcus pneumoniae	
- AD SEQUENCE: 6	
	46
81 -001 - LEMGTH: 06	
A2 RIZIZ - TYPE: DUA	
+ 7 - 1. Tr 08. MANI COI: Unknown	
+{ -1(FEATURE:	
ed dile OTHER MEDREATION: streptococcus	
ee Falor Seguence: 7	
	26
- Madyartany dissinances ageags - M. S.D.C. S.E.O. ID No. 3	69
91211 - LEUGTE: 40	
(t.) (12 × 133 E; 133)	
44 - MIB - OFWANISM: Streptocopous equi	
- Mark Come Ward And Control of C	
	46
- 97 George gegen addycz gryced gryg gagydae cedagedayeca cedageg - 99 ECTO - SEQ III No: 9	40
100 KC11 - LENGTH: 46	
100 COLO TYPE: DUA	
191 - 117 - ORGANISM: Streptococcus mutans	
104 - 0400 - SEQUENCE: 9	
105 gotttgogtg asungggtos ggggaggaat coagoagooc taagog	46
100 gooddyng ammygydda gyggaggaad doagoagdoc daagog 107 -010 × 380 lb NO: 10	4 0
100 - 1110 LENGTH: 41	
100 - 1211 - ERROGER. 41 100 - 1212 - TYPE: DMA	
110 K2130 OFGANISM: Unknown	
110 ALIGA GESANISM: OHKHOWN 111 ALIGA FEATURE:	
115 - CUSS OTHER INFORMATION: streptococcus	
115 H400H SEQUENCE: 10	1.1
116 ttgogtgsag gggtcagggg aggaatocag cagootaago g	41
118 - 1100 - SEQ IF No: 11	
119 (C111) LENGTH: 64	
110 -CITC-TYPE: DW	
171 - 0.130 OEGANIOM: Klebsiella pneumoniae	
173 -: 400: SEQUENCE: 11	.2.0
1.4 oncasegora otorgittae caggicaggi ceggaaggaa geagecagag cagaegaegi	
115 grat	64
1.7 -1.10- SEQ IL NO: 12	
118 -:111: LENGTH: 64	

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PATENT APPLICATION: US/09/766,880B

DATE: 03/21/2003
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Input Set : A:\110163-132US2.ST25.txt Output Set: N:\CRF4\03212003\I766880B.raw

100	20105 00000. 1017	
	COLOS TYPE: DNA	
	OFGANIUM: Unknown	
	TARR SHIPPER CONTROL STURBING W	
	CONTROL OF THE TAXABLE TO THE TAXABL	
1	oder - Prouence: 12	60
	: Elegona chemptimae daggioaggi beggaaggaa geagebaagg bagaigabgb	64
1.5		04
	.1.+CE, IC M: 13	
	11 - 18MGTF: (4	
	1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1	
14	-, [··· OF GAN].M: Escherichia coli	
144	-40 FF (UENCE: 1)	
145	ng nabonota ciciqtitad caggicaqgi doggaaggaa gcagocaagg dagaigadgo	60
146		64
	[LIC DEG ID NO: 34	
	11 - MEMGTH: 16	
	ANIO - TYPE: DIA	
	: F-AN.UM: Eucherichia coli	
	- 400 - SFQUENCE: 14	
	araggtragg tooggaagga agcage	26
	111 SEN IN NO: 15	
	[11] - 1FHSTF: -44	
	· Li · TYEE: DWA	
	- 19 - ANIUM: Fubudomonas aeruginosa	
	- 400 - Thatter 1	
1.	eghaschatt achogthaac otggtbaggt ooggaaggaa goagobacag ogggaabato	60
		64
$\frac{163}{163}$	·	
	+ 210 + 3EQ TF M0: 16	
	+L11 LENGTH: 04	
	+Clist TYFE: DVA	
	+U13+ OFGANISM: Excillus cereus	
170	+400. SEQUENCE: 16	ബ
	egraucqqqqa cocatgaaco ttgtcagqto oggaaggaag cagcaataag oggtottoto	64
	atut.	1,7 12
	+210	
	#211 / LENGTH: 44	
	- III TYFE: EMA	
	+213 + OPGANISM: Artificial Sequence	
179	FILTO FRATURE:	
	-173 - OTHER INFORMATION: oligonucleotide probe	
	+400 + MEQUEUCE: 17	4.4
	namaangnat annygtiagg tooggaayga agnagbagng gtgt	44
	FLIAF 7EQ II NO: 18	
	FLITT DENGTH: 30	
	+ Mil + TYPE: DNA	
1 - :-	Fully OFGANISM: Artificial Sequence	
	-Contracts:	
1 -1	+ N.C) + OTHER INFORMATION: oligonucleotide probe	
	+40] + SEQUENCE: 13	

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Input Set : A:\110163-132US2.ST25.txt Output Set: N:\CRF4\03212003\I766880B.raw

194 aggcccggga acgtattcac	20
196 KM10: SEQ ID NO: 19	
197 -:::1: LENGTH: 22	
198 -:::2: TYPE: DNA	
199 -01125 CRCANISM: Artificial Sequence	
201 HL2CH FEATURE:	
202 022 OTHER INFORMATION: oligonucleotide probe	
204 3400 SEQUENCE: 19	6.0
10% onthougheta etgegoadad gg	22
1107 (210) SEQ ID NO: 20	
HOA K211 LENGTE: 41	
AOB CAR TYPE: DNA	
M10 KM10 ORGANISM: Artificial Sequence	
MID CMD - FEATURE:	
U13 <123 - OTHER INFORMATION: oligonucleotide probe	
:15 <4.)J - SEQUENCE: 20	41
216 torgggeest tgeataagtg agtocagges tteettegts g	41

VERIFICATION SUMMARY

VERIFICATION SUMMARYDATE: 03/21/2003PATENT APPLICATION: US/09/766,880BTIME: 13:38:07

Imput Set : A:\110163-132Us2.sT25.txt Output Set: N:\CRF4\03212003\I766880B.raw